

SEQUENCE LISTING

<110> Ruben et al.

<120> Neutrokin-Alpha Binding Proteins And Methods based Thereon

<130> PF524C1

<140> Unassigned

<141> 2002-03-01

<150> 09/533,822

<151> 2000-03-24

<150> 60/188,208

<151> 2000-03-10

<150> 60/126,599

<151> 1999-03-26

<160> 5

<170> PatentIn Ver. 2.1

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<211> 882

<212> DNA

<213> Homo sapiens

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cag gag gag cgc ttt cca cag ggc ctg tgg acg ggg gtg gct atg aga	96
Gln Glu Glu Arg Phe Pro Gln Gly Leu Trp Thr Gly Val Ala Met Arg	
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Ser Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly Thr Cys Met	
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Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg Thr Cys Ala Ala	
50 55 60	
ttc tgc agg tca ctc agc tgc cgc aag gag caa ggc aag ttc tat gac	240
Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln Gly Lys Phe Tyr Asp	
65 70 75 80	
cat ctc ctg agg gac tgc atc agc tgt gcc tcc atc tgt gga cag cac	288
His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser Ile Cys Gly Gln His	

85										90										95										
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Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser Pro Val																														
100	105	110																												
aac ctt cca cca gag ctc agg aga cag cgg agt gga gaa gtt gaa aac	384																													
Asn Leu Pro Pro Glu Leu Arg Arg Gln Arg Ser Gly Glu Val Glu Asn																														
115	120	125																												
aat tca gac aac tgc gga agg tac caa gga ttg gag cac aga ggc tca	432																													
Asn Ser Asp Asn Ser Gly Arg Tyr Gln Gly Leu Glu His Arg Gly Ser																														
130	135	140																												
gaa gca agt cca gct ctc ccg ggg ctg aag ctg agt gca gat cag gtg	480																													
Glu Ala Ser Pro Ala Leu Pro Gly Leu Lys Leu Ser Ala Asp Gln Val																														
145	150	155	160																											
gcc ctg gtc tac agc acg ctg ggg ctc tgc ctg tgt gcc gtc ctc tgc	528																													
Ala Leu Val Tyr Ser Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys																														
165	170	175																												
tgc ttc ctg gtg gcg gtg gcc tgc ttc ctc aag aag agg ggg gat ccc	576																													
Cys Phe Leu Val Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro																														
180	185	190																												
tgc tcc tgc cag ccc cgc tca agg ccc cgt caa agt ccg gcc aag tct	624																													
Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser																														
195	200	205																												
tcc cag gat cac gcg atg gaa gcc ggc agc cct gtg agc aca tcc ccc	672																													
Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro																														
210	215	220																												
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Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro																														
225	230	235	240																											
acg cag gag agc gca gtc acg cct ggg acc ccc gac ccc act tgt gct	768																													
Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala																														
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Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro Cys Pro																														
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cac atc cca gac agt ggc ctt ggc att gtg tgt gtg cct gcc cag gag	864																													
His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro Ala Gln Glu																														
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<212> DNA

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atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaaac aactacaaga 540
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acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc 660
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<212> PRT

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<221> SITE

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<223> Xaa equal any amino acid

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<222> (9)

<223> Xaa equal any amino acid

<400> 4

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<210> 5

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<223> Xaa equal is any amino acid

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Ala Leu Xaa Asn Asp Glu Gly Ser Gly

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